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Muscle and serum metabolomics for different chicken breeds under commercial conditions by GC–MS

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Abstract

Globally, village chicken is popular and is known as a premium meat with a higher price. Food fraud can occur by selling other chicken breeds at a premium price in local markets. This study aimed to distinguish local village chicken from other chicken breeds available in the market, namely, colored broiler (Hubbard), broiler (Cobb), and spent laying hen (Dekalb) in pectoralis major and serum under commercial conditions using an untargeted metabolomics approach. Both pectoralis major and serum were analyzed using gas chromatography–mass spectrometry (GC–MS). The principal component analysis (PCA) results distinguished four different chicken breeds into three main groups for pectoralis major and serum. A total of 30 and 40 characteristic metabolites were identified for pectoralis major and serum, respectively. The four chicken breeds were characterized by the abundance of metabolites such as amino acids (L–glutamic acid, L–threonine, L–serine, L–leucine), organic acids (L–lactic acid, succinic acid, 3–hydroxybutyric acid), sugars (D–allose, D–glucose), sugar alcohols (myo–inositol), and fatty acids (linoleic acid). Our results suggest that an untargeted metabolomics approach using GC–MS and PCA could discriminate chicken breeds for pectoralis major and serum under commercial conditions. In this study, village chicken could only be distinguished from colored broiler (Hubbard) by serum samples. © 2021 by the authors. Licensee MDPI, Basel, Switzerland.

Author Keywords

Biomarkers; Chicken; GC–MS; Metabolomics; PCA; Pectoralis major; Serum

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